配列表

SEQUENCE LISTING

<110> ASANO Shinichiro et al.

<120> Protein Having Insecticidal Activity, DNA Coding Said Protein, Pest Control Agent and Pest Control Method

<130> BOF-3887PCT

<150> JP 2000-236140

<151> 2000-08-03

<160> 3

⟨210⟩ 1

<211> 1167

<212> PRT

<213 Bacillus thuringiensis

<400> 1

Met Ser Pro Asn Asn Gln Asn Glu Tyr Glu Ile Leu Asp Ala Ser Ser

1 5 10 15

Ser Thr Ser Val Ser Asp Asn Ser Val Arg Tyr Pro Leu Ala Asn Asp 20 25 30

Gln Thr Thr Leu Gln Asn Met Asn Tyr Lys Asp Tyr Leu Arg Met
35 40 45

Ser Glu Gly Glu Asn Pro Glu Leu Phe Gly Asn Pro Glu Thr Phe Ile 50 55 60

Ser Ser Ser Thr Val Gln Thr Gly Ile Gly Ile Val Gly Gln Val Leu
65 70 75 80

Gly Ala Leu Gly Val Pro Phe Ala Gly Gln Ile Ala Ser Phe Tyr Ser 85 90 95

- Phe Ile Val Gly Gln Leu Trp Pro Ser Ser Thr Val Ser Val Trp Glu 100 105 110
- Met Ile Met Lys Gln Val Glu Asp Leu Ile Asp Gln Lys Ile Thr Asp 115 120 125
- Ser Val Arg Lys Thr Ala Leu Ala Gly Leu Gln Gly Leu Gly Asp Gly 130 135 140
- Leu Asp Val Tyr Gln Lys Ser Leu Lys Asn Trp Leu Glu Asn Arg Asn 145 150 155 160
- Asp Thr Arg Ala Arg Ser Val Val Val Thr Gln Tyr Ile Ala Leu Glu 165 170 175
- Leu Asp Phe Val Ala Lys Ile Pro Ser Phe Ala Ile Ser Gly Gln Glu 180 185 190
- Val Pro Leu Leu Ser Val Tyr Ala Gln Ala Ala Asn Leu His Leu Leu 195 200 205
- Leu Leu Arg Asp Ala Ser Ile Phe Gly Ala Glu Trp Gly Phe Thr Pro 210 215 220
- Gly Glu Ile Ser Thr Phe Tyr Asp Arg Gln Val Thr Arg Thr Ala Gln 225 230 235 240
- Tyr Ser Asp Tyr Cys Val Lys Trp Tyr Asn Thr Gly Leu Asp Lys Leu 245 250 255
- Lys Gly Thr Asn Ala Ala Ser Trp Leu Lys Tyr His Gln Phe Arg Arg 260 265 270
- Glu Met Thr Leu Leu Val Leu Asp Leu Val Ala Leu Phe Pro Asn Tyr 275 280 285
- Asp Thr Arg Thr Tyr Pro Ile Glu Thr Thr Ala Gln Leu Thr Arg Glu

	290					295					300)			
Val 305	Tyr	Thr	Asp	Pro	11e 310	Va l	Phe	Asn	Arg	Glu 315		Ser	Gly	Gly	Phe 320
Cys	Arg	Arg	Trp	Ser 325	Leu	Asn	Ser	Asp	11e 330		Phe	Ser	Glu	Val 335	
Ser	Ala	Val	Ile 340	Arg	Ser	Pro	His	Leu 345	Phe	Asp	Ile	Leu	Ser 350	Glu	Ile
Glu	Phe	Tyr 355	Thr	Thr	Arg	Ala	Gly 360	Leu	Pro	Leu	Asn	As n 365	Thr	Glu	Tyr
Leu	Glu 370	Tyr	Trp	Val	Gly	His 375	Ser	He	Lys	Tyr	Lys 380	Asn	Thr	Asn	Ala
Ser 385	Ser	Ala	Leu	Glu	Arg 390	Asn	Tyr	Gly	Thr	Ile 395	Thr	Ser	Asn	Lys	Ile 400
Lys	Tyr	Tyr	Asp	Leu 405	Ala	Asn	Lys	Asp	11e 410	Phe	Gln	Val	Arg	Ser 415	Leu
Gly	Ala	Asp	Leu 420	Ala	Asn	Tyr	Tyr	Ala 425	Gln	Vàl	Tyr	Gly	Val 430	Pro	Tyr
Ala	Ser	Phe 435	Thr	Leu	Leu	Asp	Lys 440	Asn	Thr	Gly	Ser	Gly 445	Ser	Val	Gly
Gly	Phe 450	Thr	Tyr	Ser	Lys	Pro 455	His	Thr	Thr	Met	GI n 460	Va-l	Cys	Thr	Gln
As n 465	Tyr	Asn	Thr	He	Asp 470	Glu	He	Pro	Pro	Glu 475	Asn	Glu	Pro	Leu	Ser 480
Arg	Gly	Tyr	Ser	His	Arg	Leu	Ser	His	Ile 490	Thr	Ser	Tyr	Ser	Phe	

1008968.040302

Lys	Asn	Ala	Ser 500	Ser	Pro	Ala	Arg	Tyr 505		Asn	Leu	Pro	Val 510		Ala
Trp	Thr	His 515	Arg	Ser	Ala	Asp	Val 520	Thr	Asn	Thr	Val	Tyr 525	Ser	Asp	Lys
lle	Thr 530	Gln	lle	Pro	Val	Val 535	Lys	Ala	His	Thr	Leu 540		Ser	Gly	Thr
Thr 545	Val	Ile	Lys	Gly	Pro 550	Gly	Phe	Thr	Gly	Gly 555	Asn	Ile	Leu	Lys	Arg 560
Thr	Ser	Ser	Gly	Pro 565	Leu	Ala	Tyr	Thr	Ser 570	Val	Ser	Val	Lys	Ser 575	
Leu	Ser	Gln	Arg 580	Tyr	Arg	Ala	Arg	Ile 585	Arg	Tyr	Ala	Ser	Thr 590	Thr	Asn
Leu	Arg	Leu 595	Phe	Val	Thr	He.	Ser 600	Gly	Thr	Arg	Ile	Tyr 605	Ser	Ile	Asn
Val	Asn 610	Lys	Thr	Met	Asn	Lys 615	Gly	Asp	Asp	Leu	Thr 620	Phe	Asn	Thr	Phe
Asp 625	Leu	Ala	Thr	Ile	Gly 630	Thr	Ala	Phe	Thr	Phe 635	Ser	Asn	Туг	Ser	Asp 640
Ser	Leu	Thr	Val	Gly 645	Ala	Asp	Ser	Phe	Ala 650	Ser	Gly	Gly	Glu	Val 655	Tyr
Val	Asp	Lys	Phe 660	Glu	Leu	He	Pro	Va I 665	Asn	Ala	Thr	Phe	Glu 670	Ala	Glu
Glu	Asp	Leu 675	Asp	Val	Ala	Lys	Lys 680	Ala	Val	Asn	Gly	Leu 685	Phe	Thr	Ser

700

Lys Lys Asp Ala Leu Gln Thr Ser Val Thr Asp Tyr Gln Val Asn Gln

695

690

- Ala Ala Asn Leu Val Glu Cys Leu Ser Asp Glu Leu Tyr Pro Asn Glu 705 710 715 720
- Lys Arg Met Leu Trp Asp Ala Val Lys Glu Ala Lys Arg Leu Val Gln
 725 730 735
- Ala Arg Asn Leu Leu Gln Asp Thr Gly Phe Asn Arg Ile Asn Gly Glu 740 745 750
- Asn Gly Trp Thr Gly Ser Thr Gly Ile Glu Val Ala Glu Gly Asp Val 755 760 765
- Leu Phe Lys Asp Arg Ser Leu Arg Leu Thr Ser Ala Arg Glu Ile Asp 770 775 780
- Thr Glu Thr Tyr Pro Thr Tyr Leu Tyr Gln Gln Ile Asp Glu Ser Leu 785 790 795 800
- Leu Lys Pro Tyr Thr Arg Tyr Lys Leu Lys Gly Phe Ile Gly Ser Ser 805 810 815
- Gln Asp Leu Glu Ile Lys Leu Ile Arg His Arg Ala Asn Gln Ile Val 820 825 830
- Lys Asn Val Pro Asp Asn Leu Leu Pro Asp Val Leu Pro Val Asn Ser 835 840 845
- Cys Gly Gly Ile Asp Arg Cys Ser Glu Gln Gln Tyr Val Asp Ala Asn 850 855 860
- Leu Ala Leu Glu Asn Asn Gly Glu Asn Gly Asn Met Ser Ser Asp Ser 865 870 875 880
- His Ala Phe Ser Phe His Ile Asp Thr Gly Glu Ile Asp Leu Asn Glu 885 890 895
- Asn Thr Gly Ile Trp Val Val Phe Lys Ile Pro Thr Thr Asn Gly Tyr

			900					905					910)	
Ala	Thr	Leu 915	Gly	Asn	Leu	Glu	Leu 920		Glu	Glu	Gly	Pro 925		Ser	Gly
Glu	Thr 930	Leu	Glu	Arg	Ala	Gln 935	Gln	Gln	Glu	Gln	G1n 940		Gln	Asp	Lys
Met 945	Ala	Arg	Lys	Arg	Gly 950	Ala	Ser	Glu	Lys	Ala 955	Tyr	Tyr	Ala	Ala	Lys 960
Gln	Ala	Ile	Asp	Arg 965	Leu	Phe	Ala	Asp	Tyr 970	Gln	Asp	Gln	Lys	Leu 975	
Ser	Gly	Val	Glu 980	Met	Ser	Asp	Met	Leu 985	Ala	Ala	Gln	Asn	Leu 990	Val	Gln
Ser	Ile	Pro 995	Tyr	Val	Tyr		Asp 1000	Ala	Leu	Pro		Ile 1005	Pro	Gly	Met
	Tyr 010	Thr	Ser	Phe		Glu 1015	Leu	Thr	Asn		Leu 1020	Gln	Gln	Ala	Trp
As n 1025		Tyr	Asp		Arg 1030	Asn	Ala	lle	Pro	Asn 1035	Gly	Asp	Phe		Asn 1040
Gly	Leu	Ser		Trp 1045	Asn	Ala	Thr		Asp 1050	Val	Asn	Val		Gln 1055	Leu
Ser	Asp		Ser 1060	Val	Leu	Val		Pro 1 06 5	Asn	Trp	Asn		Gln 1070	Val	Ser
Gln		Phe 1075	Thr	Val	Gln		Asn 1080	Tyr	Arg	Tyr		Leu 1085	Arg	Val	Thr
Ala	Arg	Lys	Glu	Gly	Val	Gly	Asp	Gly	Tyr	Val	Ile	Ile	Arg	Asp	Gly

Ala Asn Gin Thr Glu Thr Leu Thr Phe Asn Ile Cys Asp Asp Asp Thr 1105 1110 1115 1120 Gly Val Leu Ser Ala Asp Gln Thr Ser Tyr Ile Thr Lys Thr Val Glu 1125 1130 1135 Phe Thr Pro Ser Thr Glu Gln Val Trp Ile Asp Met Ser Glu Thr Glu 1140 1145 1150 Gly Val Phe Asn Ile Glu Ser Val Glu Leu Val Leu Glu Glu Glu 1155 1160 1165 <210> 2 <211> 3504 <212> DNA <213 Bacillus thuringiensis <220> <221> CDS <222> (1).. (3501) <400> 2 atg agt cca aat aat caa aat gaa tat gaa att cta gat gct tca tca 48 Met Ser Pro Asn Asn Gln Asn Glu Tyr Glu Ile Leu Asp Ala Ser Ser 1 5 10 15 tot act tot gta too gat aat tot gtt aga tac cot tta gca aac gat Ser Thr Ser Val Ser Asp Asn Ser Val Arg Tyr Pro Leu Ala Asn Asp 20 25 30 caa acg acc aca tta caa aac atg aac tat aaa gat tat ctg aga atg 144 Gln Thr Thr Leu Gln Asn Met Asn Tyr Lys Asp Tyr Leu Arg Met

7/17

45

192

40

tot gag gga gag aat oot gaa tta tti gga aat oog gag acg iti att

35

Ser	Glu 50	Gly	Glu	Asn	Pro	Glu 55		Phe	Gly	Asn	Pro 60		Thr	Phe	Ile	
								att			-			-	_	240
	Ser	Ser	Thr	Val		Thr	Gly	He	Gly			Gly	Gin	Val		
65					70					75					80	
ggg	gct	tta	ggg	gtt	cca	ttt	gc t	gga	cag	ata	gct	agt	ttt	tat	agt	288
Gly	Ala	Leu	Gly	Val	Pro	Phe	Ala	Gly	Gln	He	Ala	Ser	Phe	Tyr	Ser	
				85					90					95		
								tca								336
Phe	He	Val		Gln	Leu	Trp	Pro	Ser	Ser	Thr	Val	Ser			Glu	
			100					105					110			
atσ	att	a to	222	caa	σtσ	o a a	σat	cta	att	σat	caa	222	ata	202	σat	384
								Leu								004
		115	2,0	01			120	204		. IO P	0111	125			Пор	
tct	gta	agg	aaa	aca	gcg	ctt	gca	gga	cta	caa	gga	tta	gga	gat	ggc	432
Ser	Val	Arg	Lys	Thr	Ala	Leu	Ala	Gly	Leu	Gln	Gly	Leu	Gly	Asp	Gly	
	130					135					140					
								aag								480
	Asp	Val	Tyr	GIn		Ser	Leu	Lys	Asn		Leu	Glu	Asn	Arg		
145					150					155					160	
gat	aca	aga	gc t	aga	agt	gtt	gtg	gtg	acc	caa	tat	ata	gc t	tta	gag	528
								Val								
				165					170					175		
ctt	gat	t t t	gtt	gc t	aaa	atc	cca	tct	ttt	gca	ata	t c t	gga	cag	gaa	576
Leu	Asp	Phe	Val	Ala	Lys	Ile	Pro	Ser	Phe	Ala	He	Ser	Gly	Gln	Glu	
			180					185					190			
								caa								624
vai	PTO		Leu	ser	vaı	ıyr		Gln	на	на	ASN		nıs	Leu	reu	
		195					200					205				

												Gly			cca Pro	672
							gat Asp				Thr					720
tac	_	_			gta	_	tgg Trp			act	ggc		_		tta Leu	768
		-					tgg Trp		Lys					Arg	_	816
							gat Asp 280									864
							gaa Glu									912
_							ttt Phe									960
			_	Ser 325			agt Ser									1008
							cac His			•				Glu		1056
gaa	ttt	tat	aca	aca	aga	gcg	ggg	ctt	ccc	ttg	aat	aat	acg	gaa	tac	1104

Glu	Phe	Tyr 355	Thr	Thr	Arg	Ala	Gly 360	Leu	Pro	Leu	Asn	Asn 365		Glu	Tyr	
ctt	gaa	tat	tgg	gta	gga	cat	tct	ata	aaa	tat	aaa	aat	acg	aat	gcc	1152
Leu	Glu	Tyr	Trp	Val	Gly	His	Ser	He	Lys	Туг	Lys	Asn	Thr	Asn	Ala	
	370					375					380					
tca	tca	gca	t t a	gaa	cgt	aat	tac	ggt	acg	att	ac t	t c t	aac	aaa	atc	1200
Ser	Ser	Ala	Leu	Glu	Arg	Asn	Tyr	Gly	Thr	He	Thr	Ser	Asn	Lys	He	
385					390					395					400	
aag	tat	tat	gat	t t a	gca	aat	aag	gat	atc	ttt	cag	gtt	cga	tca	t t a	1248
Lys	Tyr	Tyr	Asp	Leu	Ala	Asn	Lys	Asp	Ile	Phe	Gln	Val	Arg	Ser	Leu	
				405					410					415		
ggg	gcg	ga t	t t a	gc t	aat	tac	tac	gca	cag	gta	tat	gga	gtt	ccg	tac	1296
Gly	Ala	Asp	Leu	Ala	Asn	Tyr	Tyr	Ala	Gln	Val	Tyr	Gly	Val	Pro	Tyr	
			420					425					430			
gc t	agt	t t t	aca	ctg	ctt	gac	aag	aat	aca	gga	tca	gga	tca	gtt	gga	1344
Ala	Ser	Phe	Thr	Leu	Leu	Asp	Lys	Asn	Thr	Gly	Ser	Gly	Ser	Val	Gly	
		435					440					445				
ggt	ttt	acg	tac	tca	aaa	cca	cat	aca	ac t	atg	caa	gta	tgt	aca	caa	1392
Gly		Thr	Tyr	Ser	Lys		His	Thr	Thr	Met		Val	Cys	Thr	Gln	
	450					455					460					
aat	tac	aat	acg	att	gat	gaa	atc	cct	cca	gag	aat	gag	cca	ctt	agt	1440
Asn	Tyr	Asn	Thr	He	Asp	Glu	Ιle	Pro	Pro	Glu	Asn	Glu	Pro	Leu	Ser	
465					470					475					480	
aga	ggg	tat	agc	cat	aga	tta	t c t	cat	atc	acc	tct	tat	t c t	t t t	tct	1488
Arg	Gly	Tyr	Ser	His	Arg	Leu	Ser	His	Ile	Thr	Ser	Tyr	Ser	Phe	Ser	
				485					490					495		
aag	aat	gc t	agt	agt	cct	gc t	aga	tat	ggc	aat	ctc	c c t	gta	ttt	gct	1536
Lys	Asn	Ala	Ser	Ser	Pro	Ala	Arg	Tyr	Gly	Asn	Leu	Pro	Val	Phe	Ala	
			500					505					510			

TOORGESTOR

t gg	aca	cat	cgg	agt	gcg	gat	gtt	aca	aat	aca	gtt	tat	tca	gat	aaa	1584
Trp	Thr	His	Arg	Ser	Ala	Asp	Val	Thr	Asn	Thr	Val	Туг	Ser	Asp	Lys	
		515					520					525				
att	ac t	cag	ata	cca	gtt	gta	aag	gca	cat	ac t	tta	gtt	tca	ggt	act	1632
He	Thr	Gln	He	Pro	Val	Val	Lys	Ala	His	Thr	Leu	Val	Ser	Gly	Thr	
	530					535					540					
act	gtt	att	aaa	ggt	cct	gga	ttt	aca	gga	ggc	aat	atc	ctt	aaa	aga	1680
Thr	Val	He	Lys	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asn	Ile	Leu	Lys	Arg	
545					550					555					560	
aca	agt	agt	ggt	ccg	tta	gc t	tat	ac t	agt	gtc	tct	gta	aaa	tca	cca	1728
Thr	Ser	Ser	Gly	Pro	Leu	Ala	Tyr	Thr	Ser	Val	Ser	Val	Lys	Ser	Pro	
				565					570					575		
tta	tca	caa	aga	tat	cgt	gca	aga	ata	cgt	tat	gc t	tct	ac t	ac t	aac	1776
Leu	Ser	Gln	Arg	Tyr	Arg	Ala	Arg	He	Arg	Tyr	Ala	Ser	Thr	Thr	Asn	
			580					585					590			
tta	cga	ctt	ttt	gta	aca	att	tct	gga	ac t	cgc	att	tac	tct	ata	aat	1824
Leu	Arg	Leu	Phe	Val	Thr	He	Ser	Gly	Thr	Arg	He	Tyr	Ser	Ile	Asn	
		595					600					605				
gtt	aat	aaa	acc	atg	aat	aaa	ggg	gat	gat	tta	aca	ttt	aat	ac a	ttt	1872
Val	Asn	Lys	Thr	Met	Asn	Lys	Gly	Asp	Asp	Leu	Thr	Phe	Asn	Thr	Phe	
	610					615					620					
gac	t t a	gca	ac t	att	ggt	ac t	gc t	ttc	aca	ttt	tca	aat	tac	tcg	gat	1920
Asp	Leu	Ala	Thr	He	Gly	Thr	Ala	Phe	Thr	Phe	Ser	Asn	Tyr	Ser	Asp	
625					630					635					640	
													•	,		
agc	tta	acg	gta	ggt	gca	gat	tct	ttt	gct	tca	gga	gga	gaa	gtt	tat	1968
_											Gly					
				645		-			650		-	•		655		
gta	gat	aag	ttc	gaa	ctt	att	CCE	gta	aat	gca	aca	tti	gaa	gca	gaa	2016
g. u	0	0		044	- • •	• •	0	G • · ·		654			Gu	J 3 4	,u	0

Val Asp Lys	Phe Glu Leu 660	Ile Pro Val 665	Asn Ala Thr F	Phe Glu Ala 670	Glu
gaa gac cta	gat gtg gca	aag aaa gca	gta aat ggc t	tg ttt acg	agt 2064
Glu Asp Leu	Asp Val Ala		Val Asn Gly I		Ser
675		680	(685	
122 222 42 1	acc tta cao	ara agt gta	acg gat tat o	aa gtg aat	caa 2112
			Thr Asp Tyr (
690		695	700		
			gat gag tta t		
			Asp Glu Leu T	yr Pro Asn	720
705	710	•	715		120
aaa cga atg	tta tgg gat	gca gtg aaa	gag gcg aaa o	ga ctt gtt	cag 2208
			Glu Ala Lys A		
	725		730	735	
			+++	ott aat gga	gaa 2256
-			ttt aat agg a Phe Asn Arg		
Ald Alg Asii	740	745	THE MAIL MIG	750	
			gag gtt gcg g		
			Glu Val Ala (Val
755		760		765	
ctg ttt aaa	gat cgt tc	g ctt cgt itg	aca agt gcg	aga gag att	gat 2352
			Thr Ser Ala		
770		775	780		
			caa caa ata		
	79 Tyr Pro In		Gln Gln Ile . 795	ASP GIU SEI	800
785	13	v			
tta aaa cca	a tat aca ag	a tat aaa cta	aaa ggt ttt	ata gga agt	agt 2448
			Lys Gly Phe		
	805		810	815	5

c	aa	gat	tta	gag	att	aaa	t t a	a t a	cgt	cat	cgg	gca	aat	caa	atc	gtc	2496
								Ile									
		-		820					825					830			
2 :	ลล	aat	gta	cca	gat	aat	ctc	ttg	cca	gat	gta	ctc	cct	gtc	aat	tct	2544
								Leu									
L,	y s	ЛЭП	835	110	715 P	11011	200	840			,		845				
			000					0.10					•				
+ .	ar f	aat	aaa	atc	ga t	cgc	tøc	agt	gag	caa	cag	tat	gta	gac	gcg	aat	2592
								Ser									
U	yз	850	UIJ	110	пор	b	855	001	0.4	0111	0111	860					
		000					000					000					
ŧ	ta	ac a	ctc	o a a	aar	aat	gga	gaa	aat	gga	aat	atg	tct	tct	gat	tcc	2640
								Glu									
	65	ліц	LCu	O i u	71311	870	0.,	0.0		0.,	875				•	880	
o	00					0.0					0.0						
c	a t	gr a	111	tet	ttc	cat	att	gat	aca	ggt	gaa	ata	gat	ttg	aat	gaa	2688
		_						Asp									
11	13	,,,,,	1110	001	885					890			-		895		
					000												
а	a t	aca	gga	att	tgg	gtc	gta	ttt	aaa	att	ccg	aca	aca	aat	gga	tac	2736
		uou															
23		Thr	Glv	He	Trp												
		Thr	Gly		Trp	, 41	,			•••				910			
	.011	Thr	Gly	11e 900	Trp	741	, - 2	••••	905								
g				900					905					910			2784
	gca	aca	cta	900 gga	aat	ctt	gaa	ttg	905 gta	gaa	gag	ggg	cca	910 ttg	tca	ggg	2784
	gca	aca	cta Leu	900 gga	aat	ctt	gaa	t t g Leu	905 gta	gaa	gag	ggg	cca	910 ttg Leu	tca	ggg	2784
	gca	aca	cta	900 gga	aat	ctt	gaa	ttg	905 gta	gaa	gag	ggg	cca Pro	910 ttg Leu	tca	ggg	2784
A	gca Ma	aca Thr	cta Leu 915	900 gga Gly	aa t As n	ctt Leu	gaa Glu	ttg Leu 920	905 gta Val	gaa Glu	gag Glu	ggg Gly	cca Pro 925	910 ttg Leu	tca Ser	ggg Gly	2784 2832
A	gca Ma	aca Thr	cta Leu 915	900 gga Gly	aat Asn cga	ctt Leu gca	gaa Glu caa	ttg Leu 920 caa	905 gta Val	gaa Glu gaa	gag Glu caa	ggg Gly caa	cca Pro 925	910 ttg Leu caa	tca Ser	ggg Gly aaa	
A	gca Ma	aca Thr aca	cta Leu 915 tta Leu	900 gga Gly	aat Asn cga	ctt Leu gca	gaa Glu caa Gln	ttg Leu 920 caa Gln	905 gta Val	gaa Glu gaa	gag Glu caa	ggg Gly caa Gln	cca Pro 925 tgg	910 ttg Leu caa	tca Ser	ggg Gly aaa	
A	gca Ma	aca Thr	cta Leu 915 tta Leu	900 gga Gly	aat Asn cga	ctt Leu gca	gaa Glu caa	ttg Leu 920 caa Gln	905 gta Val	gaa Glu gaa	gag Glu caa	ggg Gly caa	cca Pro 925 tgg	910 ttg Leu caa	tca Ser	ggg Gly aaa	
A 8	gca Ala gaa Glu	aca Thr aca Thr 930	cta Leu 915 tta Leu	gga Gly gaa Glu	aat Asn cga Arg	ctt Leu gca Ala	gaa Glu caa Gln 935	tig Leu 920 caa Gln	905 gta Val caa Gln	gaa Glu gaa Glu	gag Glu caa Gln	ggg Gly caa Gln 940	cca Pro 925 tgg Trp	910 ttg Leu caa Gln	tca Ser gac Asp	ggg Gly aaa Lys	
A 8 ()	gca Ma gaa Glu	aca Thr aca Thr 930	cta Leu 915 tta Leu	900 gga Gly gaa Glu	aat Asn cga Arg	ctt Leu gca Ala	gaa Glu caa Gln 935	ttg Leu 920 caa Gln	905 gta Val caa Gln	gaa Glu gaa Glu aaa	gag Glu caa Gln	ggg Gly caa Gln 940	cca Pro 925 tgg Trp	910 ttg Leu caa Gln gca	tca Ser gac Asp	ggg Gly aaa Lys	2832
A E C	gca Ma Glu Met	aca Thr aca Thr 930 gca	cta Leu 915 tta Leu	900 gga Gly gaa Glu	aat Asn cga Arg	ctt Leu gca Ala ggg	gaa Glu caa Gln 935 gca	tig Leu 920 caa Gln	905 gta Val caa Gln	gaa Glu gaa Glu aaa	gag Glu caa Gln	ggg Gly caa Gln 940 tat Tyr	cca Pro 925 tgg Trp	910 ttg Leu caa Gln gca	tca Ser gac Asp	ggg Gly aaa Lys	2832
A E C	gca Ma gaa Glu	aca Thr aca Thr 930 gca	cta Leu 915 tta Leu	900 gga Gly gaa Glu	aat Asn cga Arg	ctt Leu gca Ala	gaa Glu caa Gln 935 gca	ttg Leu 920 caa Gln	905 gta Val caa Gln	gaa Glu gaa Glu aaa	gag Glu caa Gln gca	ggg Gly caa Gln 940 tat Tyr	cca Pro 925 tgg Trp	910 ttg Leu caa Gln gca	tca Ser gac Asp	ggg Gly aaa Lys aag Lys	2832
A 8 () () () () () () () () () () () () ()	gca Met 945	aca Thr aca Thr 930 gca Ala	cta Leu 915 tta Leu Arg	gga Gly gaa Glu aaa	aat Asn cga Arg	ctt Leu gca Ala ggg Gly 950	gaa Glu caa Gln 935 gca Ala	ttg Leu 920 caa Gln tca Ser	905 gta Val caa Gln gaa Glu	gaa Glu gaa Glu aaa Lys	gag Glu caa Gln gca Ala 955	ggg Gly caa Gln 940 tat Tyr	cca Pro 925 tgg Trp tat Tyr	910 ttg Leu caa Gln gca Ala	tca Ser gac Asp gca Ala	ggg Gly aaa Lys aag Lys	2832

Gln Ala Ile Asp	Arg Leu Phe A 965	la Asp Tyr Gln / 970	Asp Gin Lys Leu Asn 975	
tct ggt gta gaa	atg tca gat a	tg ttg gca gcc	caa aac ctt gta cag	2976
Ser Gly Val Glu	ı Met Ser Asp Mo	et Leu Ala Ala (Gln Asn Leu Val Gln	
980)	985	990	
		-4 440 000		2094
			gaa atc cct gga atg Glu Ile Pro Gly Met	3024
995	10		1005	
aac tat acg agt	ttt aca gag t	ta aca aat aga d	ctc caa caa gca tgg	3072
Asn Tyr Thr Ser	Phe Thr Glu Lo	eu Thr Asn Arg I	Leu Gln Gln Ala Trp	
1010	1015	1	020	
		nt oto aco ont o	ran ant tit aan not	3120
			gga gat tit cga aat Gly Asp Phe Arg Asn	
1025	1030	1035	1040	
1020				
gga tta agt gat	t tgg aat gca a	ca tca gat gtg a	aat gtg caa caa cta	3168
Gly Leu Ser Asp	Trp Asn Ala T	hr Ser Asp Val A	Asn Val Gln Gln Leu	
	1045	1050	1055	
aga gat aga tet	t atc ctt atc a	tt cca aac too s	aat tet caa gtg tea	3216
			Asn Ser Gln Val Ser	
1060		1065	1070	
caa caa ttt aca	a git caa ccg a	at tat aga tat g	gtg tta cgt gtc aca	3264
Gln Gln Phe Thi	r Val Gln Pro A	sn Tyr Arg Tyr '	Val Leu Arg Val Thr	
1075	10	80	1085	
~~~ ^~~ ^~~ ^~~	n ago ato ago a	ac aga tat ata	atc atc cgt gat ggt	3312
			lle lle Arg Asp Gly	
1090	1095		100	
<del>-</del>				
gcg aat cag ac	a gaa aca ctc a	ca ttt aat ata	tgt gat gat gat aca	3360
Ala Asn Gln Th	r Glu Thr Leu T	hr Phe Asn Ile	Cys Asp Asp Asp Thi	•
1105	1110	1115	112	0

ggt git tia ict gct gat caa act agc tat aic aca aaa aca gig gaa 3408 Gly Val Leu Ser Ala Asp Gln Thr Ser Tyr Ile Thr Lys Thr Val Glu 1125 1130 1135

ttc act cca tct aca gag caa gtt tgg att gac atg agt gag acc gaa 3456 Phe Thr Pro Ser Thr Glu Gln Val Trp Ile Asp Met Ser Glu Thr Glu 1140 1145 1150

ggt gta ttc aac ata gaa agt gta gaa ctc gtg tta gaa gaa gag taa 3504 Gly Val Phe Asn Ile Glu Ser Val Glu Leu Val Leu Glu Glu Glu 1155 1160 1165

<210> 3

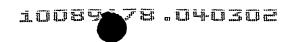
<211> 3690

<212> DNA

<213 > Bacillus thuringiensis

## <400> 3

gaattetaat gacacagtag aatattttta aaataaagat ggaagggggg atatgaaaaa 60 tataatcaca agagtcatac aaaaagatgg ttatgttaaa acaaaaaaat cctgtaggaa 120 taagggttta aaagcaatcg titgaaaaga tagttatatt aaattgtatg tataggggga 180 aaaaagatga gtccaaataa tcaaaatgaa tatgaaattc tagatgcttc atcatctact 240 tetgtateeg ataattetgt tagataeect ttageaaacg ateaaaegae cacattacaa 300 aacatgaact ataaagatta tetgagaatg tetgagggag agaateetga attattigga 360 aatccggaga cgtttattag ttcatctacg gttcaaactg gaattggcat tgttggtcaa 420 gtactggggg cittaggggt tccatttgct ggacagatag ctagtittta tagtitcatt 480 gicggicaat taiggccaic aagiaccgig agigtaiggg aaaigatiai gaaacaagig 540 gaagatetaa tigateaaaa aataacagat tetgtaagga aaacageget tgeaggaeta 600 caaggattag gagatggctt agacgtatat cagaaatcac ttaagaattg gctggaaaat 660 cgtaatgata caagagctag aagtgttgtg gtgacccaat atatagcttt agagcttgat 720 ttigtigcta aaatcccatc ttitigcaata tciggacagg aagtaccatt attatcagig 780 tatgcacaag cagcgaatti acattigcia tiattacgag atgcticcat tittiggagca 840 gagtggggat tcacaccagg agaaatttcc acattttatg atcgtcaggt gacacgtacc 900 gcccaatact cggattatig tgtaaagtgg tataacactg gcttagataa attaaaaggt 960 acgaatgctg caagttggct gaagtatcac caattccgaa gagaaatgac attactggta 1020 ttagatttag tagcgttatt tccaaactat gacacacgta cgtatccaat cgaaacaacg 1080 gcccaactta cacgggaagt gtatacagat ccaatagtat ttaacagaga aacaagtggt 1140 ggattitgta ggcgttggtc acttaacagt gatattictt titcagaagt cgaaagcgct 1200 gtaattcgtt caccacacct atttgatata ctcagtgaaa tagaatttta tacaacaaga 1260 gcggggcttc ccttgaataa tacggaatac cttgaatatt gggtaggaca ttctataaaa 1320 tataaaaata cgaatgcctc atcagcatta gaacgtaatt acggtacgat tacttctaac 1380 aaaatcaagt attatgattt agcaaataag gatatctttc aggttcgatc attaggggcg 1440 gatttagcta attactacgc acaggtatat ggagttccgt acgctagttt tacactgctt 1500 gacaagaata caggatcagg atcagttgga ggttttacgt actcaaaacc acatacaact 1560 atgcaagtat gtacacaaaa ttacaatacg attgatgaaa tccctccaga gaatgagcca 1620 citagtagag ggtatagcca tagattatct catatcacct citattcttt tictaagaat 1680 gctagtagtc ctgctagata tggcaatctc cctgtatttg cttggacaca tcggagtgcg 1740 gatgitacaa atacagitta ticagataaa attacicaga taccagitgi aaaggcacai 1800 actitagtit caggiactae igitatiaaa ggicciggat tiacaggagg caataiccit 1860 aaaagaacaa gtagtggtcc gttagcttat actagtgtct ctgtaaaatc accattatca 1920 caaagatatc gigcaagaat acgitaigci iciaciacia acitacgaci iiitgiaaca 1980 atticiggaa cicgcattta cictataaat gitaataaaa ccatgaataa aggggatgat 2040 ttaacattta atacatttga cttagcaact attggtactg ctttcacatt ttcaaattac 2100 teggataget taaeggtagg tgeagattet titgetteag gaggagaagt tiatgtagat 2160 aagticgaac ttattccggt aaatgcaaca tttgaagcag aagaagacct agatgtggca 2220 aagaaagcag taaatggctt gtttacgagt aaaaaagatg ccttacagac aagtgtaacg 2280 gattatcaag tgaatcaagc ggcaaactta gtagaatgcc tatccgatga gttataccca 2340 aatgaaaaac gaatgttatg ggatgcagtg aaagaggcga aacgacttgt tcaggcacgt 2400 aacttactcc aagatacagg cittaatagg attaatggag aaaacggatg gacgggaagt 2460 acgggaatcg aggttgcgga aggagatgit ctgtttaaag atcgttcgct tcgtttgaca 2520 agtgcgagag agattgatac agaaacatat ccaacgtatc tctatcaaca aatagatgaa 2580 tcacttttaa aaccatatac aagatataaa ctaaaaggtt ttataggaag tagtcaagat 2640 ttagagatta aattaatacg tcatcgggca aatcaaatcg tcaaaaatgt accagataat 2700 cictigccag aigiactccc igicaatici igiggiggga icgaicgcig cagigagcaa 2760 cagtatgtag acgcgaattt agcactcgaa aacaatggag aaaatggaaa tatgtcttct 2820 gattcccatg cattttcttt ccatattgat acaggtgaaa tagatttgaa tgaaaataca 2880 ggaatttggg tcgtatttaa aattccgaca acaaatggat acgcaacact aggaaatctt 2940 gaatiggiag aagaggggcc atigicaggg gaaacattag aacgagcaca acaacaagaa 3000 caacaatggc aagacaaaat ggcaagaaaa cgtggggcat cagaaaaagc atattatgca 3060 gcaaagcaag ccattgatcg titaticgca gattatcaag accaaaaact taatictggt 3120 gtagaaatgt cagatatgtt ggcagcccaa aacctigtac agtccattcc ttacgtatat 3180 aatgatgcgt taccagaaat ccctggaatg aactatacga gitttacaga gitaacaaat 3240



agactccaac aagcatggaa titgtatgat citcgaaatg ciataccaaa iggagatiii 3300 cgaaatggat taagtgatig gaatgcaaca icagatgtga atgtgcaaca actaagcgat 3360 acatctgtcc tigicaticc aaactggaat tcicaagtgi cacaacaati tacagticaa 3420 ccgaattata gatatgtgit acgtgtcaca gcgagaaaag agggagtagg agacggatat 3480 gigatcatcc gigatggtgc gaatcagaca gaaacactca catttaatat atgtgatgat 3540 gatacaggtg tittatctgc igatcaact agctatatca caaaaacagt ggaattcact 3600 ccatctacag agcaagtitg gattgacatg agtaggaccg aaggtgat caacatagaa 3660 agtgtagaac tcgtgitaga agaagagtaa